

University of Groningen

Bioinformatics for mass spectrometry. Novel statistical algorithms

Dijkstra, Martijn

IMPORTANT NOTE: You are advised to consult the publisher's version (publisher's PDF) if you wish to cite from it. Please check the document version below.

Document Version

Publisher's PDF, also known as Version of record

Publication date:

2008

[Link to publication in University of Groningen/UMCG research database](#)

Citation for published version (APA):

Dijkstra, M. (2008). *Bioinformatics for mass spectrometry. Novel statistical algorithms*. s.n.

Copyright

Other than for strictly personal use, it is not permitted to download or to forward/distribute the text or part of it without the consent of the author(s) and/or copyright holder(s), unless the work is under an open content license (like Creative Commons).

The publication may also be distributed here under the terms of Article 25fa of the Dutch Copyright Act, indicated by the "Taverne" license. More information can be found on the University of Groningen website: <https://www.rug.nl/library/open-access/self-archiving-pure/taverne-amendment>.

Take-down policy

If you believe that this document breaches copyright please contact us providing details, and we will remove access to the work immediately and investigate your claim.

Downloaded from the University of Groningen/UMCG research database (Pure): <http://www.rug.nl/research/portal>. For technical reasons the number of authors shown on this cover page is limited to 10 maximum.

Publications

- Fu J, **Dijkstra M**, Keurentjes JJB, Koorneef M, Breitling R and Jansen RC, *Systems biology through system genetics*, **in preparation**
- **Dijkstra M** and Jansen RC, *Advanced deconvolution analysis of complex mass spectra*, **submitted**
- Roelofsen H*, Alvarez-Llamas G*, **Dijkstra M***, Breitling R, Havenga K, Bijzet J, Zandbergen W, De Vries M, Ploeg R and Vonk RJ (2007) *Analyses of intricate kinetics of the serum proteome during and after colon surgery by protein expression time series*, **Proteomics** 7: 3219–3228
*Equal contribution doi:10.1002/pmic.200601047
- **Dijkstra M**, Vonk RJ and Jansen RC (2007) *SELDI-TOF mass spectra: a view on sources of variation*, **Journal of Chromatography B** 847: 12–23
doi:10.1016/j.jchromb.2006.11.004
- **Dijkstra M**, Roelofsen H, Vonk RJ and Jansen RC (2006) *Peak quantification in surface-enhanced laser desorption/ionization by using mixture models*, **Proteomics** 6: 5106–5116
doi:10.1002/pmic.200600181

